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IN THE CLAIMS:

Please cancel claims 30, 32, 33, 36-39, 40-52, and 55-61 without prejudice or disclaimer.

The pending claims are listed in the attached:

Appendix A1: Pending claims (clean copy).

Changes in the pending claims relative to the last version of record are reflected \boxtimes in:

Appendix A2: Changes to claims (redline).

Please enter any new claims or changes reflected in Appendices A1 and A2.

REMARKS

Claims 27-61 were pending. Claims 27-61 were subject to a seven (7) way Restriction Election/Restriction 7 27 39 7 53, c, 4 and 6 (
secute Group I (claims ~ Requirement. In addition, Claims 27-30, 53, and 54 have been amended. Support for the amendments is apparent. Thus, no new matter is added.

Applicants hereby elect to prosecute Group I (claims 27-30, 34, 35 53, and 54), without traverse. While the election of Group I is without traverse, Applicants reserve the right to traverse restriction requirements between the other asserted claim groups. Applicants provisionally elect SEQ ID NO:2, but will seek to keep the highly related sequences together.

FEE DEFICIENCY

If an extension of time is deemed required for consideration of this paper, please consider 冈 this paper to comprise a petition for such an extension of time; The Commissioner is hereby authorized to charge the fee for any such extension to Deposit Account No. 50-0258.

and/or

If any additional fee is required for consideration of this paper, please charge Account X No. 50-0258.

Closing Remarks

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Entry of the Response & Amendment and allowance of the pending claims are respectfully requested.

Respectfully submitted,

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Appendix A2: Changes to claims (redline).

- 27. An isolated polypeptide comprising a member selected from the group consisting of
 - (a) an amino acid sequence which has at least 90% identity to matching SEQ ID NO:22, SEQ ID NO:4, SEQ ID NO:6, or SEQ ID NO:8;
 - (b) an immunogenic <u>polypeptide comprising a</u> fragment of the amino acid sequence of (a), wherein the immunogenic fragment is at least 90% identical to an aligned contiguous segment of SEQ ID NO:2; and(c) an immunogenic fragment of the 15 amino acid sequence of (a) acids that matches an aligned contiguous segment of SEQ ID NO:2 with no more than five single amino acid substitutions, deletions 2, SEQ ID NO:4, SEQ ID NO:6, or additions SEQ ID NO:8;

wherein the isolated polypeptide, when administered to a subject in a suitable composition which can include an adjuvant, or a suitable carrier coupled to the polypeptide, induces an <u>antibody or <u>T-cell</u> immune response that recognizes to a polypeptide having the sequence of SEQ ID NO:2-2. SEQ ID NO:4, SEQ ID NO:6, or SEQ ID NO:8;</u>

- The isolated polypeptide of claim 27, wherein the immune response recognizes an antigen having molecular weight 17-18kD (by SDS PAGE) in *Moraxella catarrhalis* strains 2904, 2905, 2906, 2907, 2908, 2909, 2910, 2911, 2912, 2913, 2926, 2931, 2956, 2960, 2969, and 2975. polypeptide is according to (a).
- 29. The isolated polypeptide of claim 27, wherein the immunogenic fragment of polypeptide is according to (b) comprises at least 15 amino acids.
- 31. The isolated polypeptide of claim 27, wherein the immunogenic fragment of (b) comprises at least 20 amino acids.
- 34. The isolated polypeptide of claim 3227, wherein the isolated polypeptide consists of the amino acid sequence of SEQ ID NO:2.2, SEQ ID NO:4, SEQ ID NO:6, or SEQ ID NO:8.
- 54. The vaccine of Claim 53, wherein the composition vaccine comprises at least one other Moraxella catarrhalis antigen.